Substitute Page 1



SPECIFICALLY TARGETED ANTIBODY AGENT, COMPOSITION, KIT & RELATED USES

BACKGROUND OF THE INVENTION

Related Patents

This application is a divisional of US Patent Application Serial No. 08/482,596, filed June 7, 1995, which has been allowed; which in turn is a divisional of USSN 08/162,402 filed December 3, 1990, now US Patent No. 5,972,337; which is a continuation-in-part of USSN 07/607,538, now US Patent No. 5,455,031.

Technical Field

invention relates to a polypeptide having This **HMFG** 46 kDalton antibody binding specificity of the polynucleotide, а and antigen, a differentiation polyribonucleotide encoding it, anti-polypeptide antibodies, methods of detecting the polypeptide and DNA and RNA encoding it, a method of imaging cells expressing the polypeptide, a method of detecting the presence of the polypeptide in a biological fluid by binding the antibody to the polypeptide, in vivo and ex vivo methods of delivering a therapeutic agent to a target cell expressing the polypeptide, a fusion protein of the polypeptide and at least one other polypeptide, labeled polynucleotides and polyribonucleotides encoding the polypeptide and a complementary DNA sequence, method of detecting RNA and DNA by hybridization with labeled probes, a method of vaccination with the polypeptide, and method of treating breast cancer with an anti-sense DNA.

WHAT IS SEEKED TO BE PATENTED AS NOVEL & UNOBVIOUS IN LETTERS PATENT OF THE UNITED STATES IS:

- 52. (Canceled)
- 53. (Canceled)
- 54. (Canceled)
- 55. (Canceled)
- 56. (Canceled)
- 57. (Canceled)
- 58. (Canceled)
- 59. (Canceled)
- 60. (Canceled)
- 61. (Canceled)
- 62. (Canceled)
- 63. (Canceled)
- 64. (Canceled)
- 65. (Canceled)
- 66. (Canceled)
- 81. (Amended) An in vivo method of imaging a cancer of epithelial origin or cells expressing a polypeptide having the antibody binding specificity of the about 46 Kd differentiation Human Milk Fat Globule (HMFG) antigen, comprising

administering to a subject suspected of being afflicted with the cancer or carrying the cells an amount of a detectably labeled or unlabeled specifically targeted antibody, comprising a monoclonal antibody selectively binding a 46 Kd MW HMFG differentiation antigen that has an antigen affinity constant about 10^{10} - 10^{5} M⁻¹, and an agent comprising a detectable label, the antibody and the agent being operatively linked to one another, under conditions effective to deliver the antibody to target cells of epithelial origin carrying at least a portion of the 46Kd MW HMFG differentiation antigen in the subject's body to form antibody-cell antigen complexes;

administering to the subject a detectably labeled agent that binds the antibody at a site other than the 46 kDalton HMFG polypeptide binding site if the antibody is unlabeled; and detecting the presence of a label in the subject's body.

- 82. (Amended) The method of claim 81, wherein the antibody is administered intravenously, intraperitoneally, intracavitarily, intra-tumor, intramuscularly, or into the lymphatic system.
- 83. (Amended) The method of claim 81, wherein the labeled agent comprises a fuorescent or radiolabeled agent.
 - 84. (Amended) The method of claim 81, wherein

the antibody comprises an unlabeled antibody; and

the labeled agent comprises a labeled anti-antibody immunoglobulin, antibody binding fragment thereof, protein A, or Protein C.

85. (Amended) The method of claim 81, further comprising upon label detection the delivery of a therapeutic agent to target cancerous cells or cells of epithelial origin by

binding a therapeutic agent to the antibody of claim 81, at a site other than its antigen binding site;

administering to the subject a therapeutically effective amount of the antibody-bound therapeutic agent under conditions effective for the antibody to deliver the agent to the target cells; and

allowing the antibody to bind to the target cells, and the therapeutic agent to exert its effect on the cells.

- 86. (Canceled)
- 87. (Canceled)
- 88. (New) The method of claim 81, wherein the labeled agent comprises a radionucleide, a fluorescent label, an enzyme or biotin.
- 89. (New) The method of claim 81, wherein the labeled agent is detected as a conjugate.

- 90. (New) The method of claim 89, wherein the antibody is conjugated to avidin, streptavidin, or a magnetic bead.
- 91. (New) The method of claim 81, wherein the antibody comprises a monoclonal antibody.
- 92. (New) The method of claim 81, wherein the antibody is provided as a composition with a non-proteolytic carrier.
- 93 (New) The method of claim 92, wherein the carrier comprises a biologically acceptable carrier.
- 94. (New) The method of claim 93, wherein the carrier comprises a pharmaceutically acceptable carrier.
- 95. (New) The method of claim 85, wherein the therapeutic agent comprises a radionucleide, an immmunotoxin, or an enzyme.
- 96. (New) The method of claim 85, wherein the antibody-therapeutic agent is delivered as a conjugate.
- 97. (New) The method of claim 96, wherein the antibody-therapeutic agent is conjugated to avidin, streptavidin, or a magnetic bead.
- 98. (New) The method of claim 85, wherein the antibody-therapeutic agent comprises a monoclonal antibody.
- 99. (New) The method of claim 85, wherein antibody-therapeutic agent is provided as a composition with a non proteolytic carrier.
- 100. (New) The method of claim 99, wherein the antibody-therapeutic agent carrier comprises a biologically acceptable carrier.
- 101. (New) The method of claim 100, wherein the antibody-therapeutic agent carrier comprises a pharmaceutically acceptable carrier.

THE 1 2005 TO TAKE

Table 1: DNA Sequence and Derived Amino Acid Sequence of BA46-1 cDNA.

| , | * | 1 | o I | * | | 20 | | * | 30 | | * | 4 | 0 | * | | 50 I | | |
|--|---|---|--|--|--|---|---|---|---|--|--|--|--|---|---|--|---|--|
| GAT TT Asp Ph | rc A | ATC Ile | CAT His | GAT Asp | GTT Val | AAT Asn | AAA Lys | AAA Lys | CAC His | AAG Lys | GAG Glu | TTT Phe | GTG Val | GGT Gly | AAC Asn | TGG Trp | AAC Asn | |
| * | 60 | | * | | 70 | * | | 80 | | * | 90 | 1 | * | 1 | .00 | * | | |
| AAA AA Lys As | AC (| GCG Ala | GTG Val | CAT His | GTC Val | AAC Asn | CTG Leu | TTT Phe | GAG Glu | ACC Thr | CCT Pro | GTG Val | GAG Glu | GCT Ala | ĊAG Gln | TAC Tyr | GTG Val | |
| 110 | • | * | 120 | | * | 130 |) | * | 14 | 10 | * | 1 | .50 | | * | 160 |) | |
| AGA T | TG ' eu ' | TAC Tyr | CCC Pro | ACG Thr | AGC Ser | TGC Cys | CAC His | ACG Yhr | GCC Ala | TGC Cys | ACT Thr | CTG Leu | CGC Arg | TTT Phe | GAG Glu | CTA Leu | CTG Leu | |
| * | | 170 | | * | 180 |) | * | 1 | 190 | , | ٠ | 200 | | * | 210 |) | * | |
| GGC TO | GT (ys (| GAG Glu | CTG Leu | AAC Asn | GGÅ Gly | TGC Cys | GCC Ala | AAT Asn | ccc Pro | CTG Leu | GGC Gly | CTG Leu | AAG Lys | AAT Asn | AAC Asn | AGC Ser | ATC Ile | |
| 220 | 0 | * | | 230 | | * | 240 |) | * | 2 | 250 | * | • | 260 | | * | 270 | |
| CCT G | AC sp | AAG Lys | CAG Gln | ATC Ile | ACG Thr | GCC Ala | TCC Ser | AGC Ser | AGC Ser | TAC Tyr | AAG Lys | ACC Thr | TGG Trp | GGC Gly | TTG Leu | CAT His | CTC Leu | |
| , | * | 2 | 80 | , | + | 290 | | * | 300 | | * | 3 | 310 | ; | + | 320 | | |
| TTC A | GC er | TGG Trp | AAC Asn | CCC Pro | TCC Ser | TAT Tyr | GCA Ala | CGG Arg | CTĠ Leu | GAC Asp | AAG Lys | CAG Gln | GGC Gly | AAC Asn | TTC Phe | AAC Asn | GCC Ala | |
| * | 330 | | * | ; | 340 | , | * | 350 | | * | 360 |) | * | ; | 370 | 1 | t | |
| TGG G | TT al | GCG Ala | GGG Gly | AGC Ser | TAC Tyr | GGT Gly | AAC Asn | GAT Asp | CAG Gln | TGG Trp | CTG Leu | CAG Gln | GTG Val | GAC Asp | ĊTG Leu | GGC Gly | TCC Ser | |
| 380 | | * | 390 | | * | 4 | 90 | * | | 410 | | * | 420 | | * | 4 | 130 | |
| TCG A Ser L | AG ys | GAG Glu | GTG Val | ACA Thr | GGC Gly | ATC Ile | ATC Ile | ACC Thr | CAG Gln | GĠG Gly | GCC Ala | CGT Arg | AAC Asn | TTT Phe | GGC Gly | TCT Ser | GTC Val | |
| * 4 | 40 | | * | 45 | 0 | * | • | 460 | * | | 470 | | * | 48 | 0 | * | | |
| CAG T Gln P | | | | | | | | | | | | | | | | | | |
| 490 | | * | 500 | * | | 51 | 0 | * | | 520 | , | * | 530 | | * | 541 | 0 | |
| GAG T Glu T | AC 'yr | CAG Gln | GAC Asp | CCC Pro | AGG Arg | ACT Thr | GGC Gly | AGC Ser | AGT Ser | AAG Lys | ATC Ile | TTC Phe | CCT Pro | GGC Gly | AAC Asn | TGĠ Trp | GAC Asp | |
| | | į | 550 | | * | 560 | | * | 57 I | 0 | * | ! | 580 I | | * | 590 I | | |
| AAC C Asn H | AC lis | TCC Ser | CAC His | AAG Lys | AAG Lys | AAC Asn | TTG Leu | TTT Phe | GAĠ Glu | ACG Thr | CCC Pro | ATC Ile | CTG Leu | GCT Ala | CGC Arg | ŤAT Tyr | GTG Val | |
| * 6 | 00 | | * | 61 | 0 | * | | 620 I | | * | 630 I | | * | 64 | 0 | * | | |
| CGC A Arg I | TC le | CTG Leu | CCT Pro | GTA Val | GCC Ala | TGG Trp | CAC His | AAC Asn | CGC Arg | ATC Ile | GĊC Ala | CTG Leu | CGC Arg | CTG Leu | GÁG Glu | CTG Leu | CTG Leu | |
| 650 I | | * | 660 | | * | 67 1 | 0 * | | | 680 I | | * | 690 I | | * | 70 I | 0 | |
| | | | | | | | | | | | | | | ~~~ | | | CCC | 703 |
| GGC T Gly C | | | | | CCT NO: | | ACC | ccc | AGG | TCT | TCC | TGC | TTT | CCA | TGG | GCC | CGC | |
| Gly C | ys TC | End TTG | (GCT | SEQ. TCT | NO: | 2) CCC | CTT | TAA | ATC | ACC | ATA | GGG | CTG | GGG | ACT | GGG | GAA | 757 811 |
| TGC C GGG G CCC A | TC SAG ACC | End TTG GGT CTC | GCT GTT CAC | SEQ. TCT CAG CTC | NO: CAG AGG TCA | 2) CCC CAG CGG | CTT CAC GCC | TAA CAC CTG | ATC CAC CCC | ACC ACA CAG | ATA GTC CCC | GGG ACC CTA | CTG CCT AGC | GGG CCC | ACT TCC GTC | GGG CTC CCC | GAA TTT TAA | |
| TGC CGGG GCCC ACCC CGGG A | TC SAG ACC CCA ATG | TTG GGT CTC GTC GAC | GCT GTT CAC CTC AGG | TCT CAG CTC ACT AAA | CAG AGG TCA GTC GGG | CCC CAG CGG CTG CAA | CTT CAC GCC TTT AGT | TAA CAC CTG TCT AGG | ATC CAC CCC TAG GCG | ACC ACA CAG GCA | ATA GTC CCC CTG | GGG ACC CTA AGG TTC | CTG CCT AGC GAT CCT | GGG CCC CCC CTG | ACT TCC GTC AGT CCT | GGG CTC CCC AGG | GAA TTT TAA TCT CGG | 811 865 |
| Gly C TGC C GGG G CCC A CCC C GGG A ACC G | YS SAG ACC CCA ATG SCC | TTG GGT CTC GTC GAC GAT TCC | GCT GTT CAC CTC AGG CCC CAT | TCT CAG CTC ACT AAA AGG GGT | CAG AGG TCA GTC GGG TGC | CCC CAG CGG CTG CAA GTG | CTT CAC GCC TTT AGT TGT AAG | TAA CAC CTG TCT AGG CTC | ATC CAC CCC TAG GCG TGT | ACC ACA CAG GCA TGT CTC | ATA GTC CCC CTG GGT TCC | GGG ACC CTA AGG TTC TAG GCC | CTG CCT AGC GAT CCT CCC CCA | GGG CCC CTG GCC CTC | ACT TCC GTC AGT CCT TCT | GGG CTC CCC AGG GTC CAC | GAA TTT TAA TCT CGG ACA TAA | 811 865 919 973 |
| Gly C GGG G CCC A CCC C GGG A ACC G TCA C CAG C | YS SAG ACC ATG SCC AT SCC SAT SCT | End TTG GGT CTC GAC GAT TCC CTT GCC | GCT GTT CAC CTC AGG CCC CAT GCC | TCT CAG CTC ACT AAA AGG GGT CGT | CAG AGG TCA GTC GGG TGC CGG CCC | CCC CAG CTG CTAA GTG CTC | CTT CAC GCC TTT AGT TGT AAG TGC | TAA CAC CTG TCT AGG CTC AAA GTC | ATC CAC CCC TAG GCG TGT GGC GGC | ACC ACA CAG GCA TGT CTC | ATA GTC CCC CTG GGT TCC GAA GGG | GGG ACC CTA AGG TTC TAG GCC GTA | CTG CCT AGC GAT CCT CCC CCA CCA | GGG CCC CTG GCC CTC GGC TGT | ACT TCC GTC AGT CCT TCT TGG GCC | GGG CTC CCC AGG GTC CAC AGA ACA | GAA TTT TAA TCT CGG ACA TAA ACT TCT | 811 865 919 973 1027 1081 |
| Gly C TGC C GGG G CCC A CCC C GGG A ACC G TCA C | CYS CTC CAC CCA CCA CCA CCT CCT CCT CCT CCT | End TTG GGT CTC GAC GAT TCC CTT GCC TTG GGG | GCT GTT CAC CTC AGG CCC CAT GCC TCC AGC | TCT CAG CTC ACT AAA AGG GGT CGT TGA GAG | NO: CAG AGG TCA GTC GGG CGG CGG CCC AGC GTC | CCC CAG CTG CTAA GTG CTC CCAA CCAA | CTT CAC GCC TTT AGT TGT AAG GAC GCG | TAA CAC CTG TCT AGG CTC AAA GTC ACT | ATC CAC CCC TAG GCG TGT GGC TCC CAG | ACC ACA CAG GCA TGT CTC CCG CCT | ATA GTC CCC CTG GGT TCC GAA GGG TGT GGG | GGG ACC CTA AGG TTC TAG GCC GTA CTC GTG | CTG CCT AGC GAT CCT CCA CCA CCT GGG GGG | GGG CCC CTG GCC CTC GGC TGT CGG TGG | ACT TCC GTC AGT CCT TCT TGG GCC TGC | GGG CTC CCC AGG GTC CAC AGA ACA CTC TAT | GAA TTT TAA TCT CGG ACA TAA ACT TCT GGG GGG | 811 865 919 973 1027 1081 1135 1189 |

Potential n-linked glycosylation sites are underlined.

Table 2: Comparison of Derived BA46-1 Amino Acid Sequence with C-terminal Human Serum Factors V and VIII

An arrow indicates Junction of C1 and C2 repeats.